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<213> Mus musculus

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35 40 45

Gly Arg Lys Arg Lys His Pro Pro Val Glu Ser Ser Asp Thr Pro Lys
50 55 60

Asp Pro Ala Val Thr Thr Lys Ser Gln Pro Met Ala Gln Asp Ser Gly
65 70 75 80

Pro Ser Asp Leu Leu Pro Asn Gly Asp Leu Glu Lys Arg Ser Glu Pro
85 90 95

Gln Pro Glu Glu Gly Ser Pro Ala Ala Gly Gln Lys Gly Gly Ala Pro
100 105 110

Ala Glu Gly Glu Gly Thr Glu Thr Pro Pro Glu Ala Ser Arg Ala Val
115 120 125

Glu Asn Gly Cys Cys Val Thr Lys Glu Gly Arg Gly Ala Ser Ala Gly
130 135 140

Glu Gly Lys Glu Gln Lys Gln Thr Asn Ile Glu Ser Met Lys Met Glu
145 150 155 160

Gly Ser Arg Gly Arg Leu Arg Gly Gly Leu Gly Trp Glu Ser Ser Leu
165 170 175

Arg Gln Arg Pro Met Pro Arg Leu Thr Phe Gln Ala Gly Asp Pro Tyr
180 185 190

Tyr Ile Ser Lys Arg Lys Arg Asp Glu Trp Leu Ala Arg Trp Lys Arg
195 200 205

Glu Ala Glu Lys Lys Ala Lys Val Ile Ala Val Met Asn Ala Val Glu
210 215 220

Glu Asn Gln Ala Ser Gly Glu Ser Gln Lys Val Glu Glu Ala Ser Pro
225 230 235 240

Pro Ala Val Gln Gln Pro Thr Asp Pro Ala Ser Pro Thr Val Ala Thr

100 110 120 130 140 150 160 170 180 190 200 210 220 230 240

245

250

255

Thr Pro Glu Pro Val Gly Gly Asp Ala Gly Asp Lys Asn Ala Thr Lys
260 265 270

Ala Ala Asp Asp Glu Pro Glu Tyr Glu Asp Gly Arg Gly Phe Gly Ile
275 280 285

Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe Ser Trp Trp Pro Gly
290 295 300

Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser Arg Ala Ala Glu Gly
305 310 315 320

Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys Phe Ser Val Val Cys
325 330 335

Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys Ser Ala Phe His Gln
340 345 350

Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys Ala Ile Tyr Glu Val
355 360 365

Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu Phe Pro Ala Cys His
370 375 380

Asp Ser Asp Glu Ser Asp Ser Gly Lys Ala Val Glu Val Gln Asn Lys
385 390 395 400

Gln Met Ile Glu Trp Ala Leu Gly Gly Phe Gln Pro Ser Gly Pro Lys
405 410 415

Gly Leu Glu Pro Pro Glu Glu Lys Asn Pro Tyr Lys Glu Val Tyr
420 425 430

Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala Tyr Ala Pro Pro Pro
435 440 445

Pro Ala Lys Lys Pro Arg Lys Ser Thr Thr Glu Lys Pro Lys Val Lys

450 455 460
Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu Val Tyr Glu Val Arg
465 470 475 480

Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile Ser Cys Gly Ser Leu
485 490 495

Asn Val Thr Leu Glu His Pro Leu Phe Ile Gly Gly Met Cys Gln Asn
500 505 510

Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln Tyr Asp Asp Asp Gly
515 520 525

Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly Arg Glu Val Leu Met
530 535 540

Cys Gly Asn Asn Asn Cys Cys Arg Cys Phe Cys Val Glu Cys Val Asp
545 550 555 560

Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala Ile Lys Glu Asp Pro
565 570 575

Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr Tyr Gly Leu Leu Arg
580 585 590

Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met Phe Phe Ala Asn Asn
595 600 605

His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr Pro Pro Val Pro Ala
610 615 620

Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala
625 630 635 640

Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile Gln Val Asp Arg Tyr
645 650 655

Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr Val Gly Met Val Arg

DRAFT VERSION 1

660

665

670

His Gln Gly Lys Ile Met Tyr Val Gly Asp Val Arg Ser Val Thr Gln
675 680 685

Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu Val Ile Gly Gly Ser
690 695 700

Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala Arg Lys Gly Leu Tyr
705 710 715 720

Glu Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr Arg Leu Leu His Asp
725 730 735

Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe Phe Trp Leu Phe Glu
740 745 750

Asn Val Val Ala Met Gly Val Ser Asp Lys Arg Asp Ile Ser Arg Phe
755 760 765

Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys Glu Val Ser Ala Ala
770 775 780

His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly Met Asn Arg Pro
785 790 795 800

Leu Ala Ser Thr Val Asn Asp Lys Leu Glu Leu Gln Glu Cys Leu Glu
805 810 815

His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg Thr Ile Thr Thr Arg
820 825 830

Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His Phe Pro Val Phe Met
835 840 845

Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu Met Glu Arg Val Phe
850 855 860

Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn Met Ser Arg Leu Ala

865

870

875

880

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Asp Thr Lys Asp Ala Pro Ser Pro Pro Val Leu Glu Ala Ile Cys Thr
35 40 45

Glu Pro Val Cys Thr Pro Glu Thr Arg Gly Arg Arg Ser Ser Ser Arg
50 55 60

Leu Ser Lys Arg Glu Val Ser Ser Leu Leu Asn Tyr Thr Gln Asp Met
65 70 75 80

Thr Gly Asp Gly Asp Arg Asp Asp Glu Val Asp Asp Gly Asn Gly Ser
85 90 95

Asp Ile Leu Met Pro Lys Leu Thr Arg Glu Thr Lys Asp Thr Arg Thr
100 105 110

Arg Ser Glu Ser Pro Ala Val Arg Thr Arg His Ser Asn Gly Thr Ser
115 120 125

Ser Leu Glu Arg Gln Arg Ala Ser Pro Arg Ile Thr Arg Gly Arg Gln
130 135 140

Gly Arg His His Val Gln Glu Tyr Pro Val Glu Phe Pro Ala Thr Arg
145 150 155 160

Ser Arg Arg Arg Arg Ala Ser Ser Ser Ala Ser Thr Pro Trp Ser Ser
165 170 175

Pro Ala Ser Val Asp Phe Met Glu Glu Val Thr Pro Lys Ser Val Ser
180 185 190

Thr Pro Ser Val Asp Leu Ser Gln Asp Gly Asp Gln Glu Gly Met Asp
195 200 205

Thr Thr Gln Val Asp Ala Glu Ser Arg Asp Gly Asp Ser Thr Glu Tyr
210 215 220

Gln Asp Asp Lys Glu Phe Gly Ile Gly Asp Leu Val Trp Gly Lys Ile
225 230 235 240

Lys Gly Phe Ser Trp Trp Pro Ala Met Val Val Ser Trp Lys Ala Thr
245 250 255

Ser Lys Arg Gln Ala Met Pro Gly Met Arg Trp Val Gln Trp Phe Gly
260 265 270

Asp Gly Lys Phe Ser Glu Ile Ser Ala Asp Lys Leu Val Ala Leu Gly
275 280 285

Leu Phe Ser Gln His Phe Asn Leu Ala Thr Phe Asn Lys Leu Val Ser
290 295 300

Tyr Arg Lys Ala Met Tyr His Thr Leu Glu Lys Ala Arg Val Arg Ala
305 310 315 320

Gly Lys Thr Phe Ser Ser Ser Pro Gly Glu Ser Leu Glu Asp Gln Leu
325 330 335

Lys Pro Met Leu Glu Trp Ala His Gly Gly Phe Lys Pro Thr Gly Ile
340 345 350

Glu Gly Leu Lys Pro Asn Lys Lys Gln Pro Val Val Asn Lys Ser Lys
355 360 365

Val Arg Arg Ser Asp Ser Arg Asn Leu Glu Pro Arg Arg Arg Glu Asn
370 375 380

Lys Ser Arg Arg Arg Thr Thr Asn Asp Ser Ala Ala Ser Glu Ser Pro
385 390 395 400

Pro Pro Lys Arg Leu Lys Thr Asn Ser Tyr Gly Gly Lys Asp Arg Gly
405 410 415

Glu Asp Glu Glu Ser Arg Glu Arg Met Ala Ser Glu Val Thr Asn Asn
420 425 430

Lys Gly Asn Leu Glu Asp Arg Cys Leu Ser Cys Gly Lys Lys Asn Pro
435 440 445

Val Ser Phe His Pro Leu Phe Glu Gly Leu Cys Gln Ser Cys Arg
450 455 460

Asp Arg Phe Leu Glu Leu Phe Tyr Met Tyr Asp Glu Asp Gly Tyr Gln
465 470 475 480

Ser Tyr Cys Thr Val Cys Cys Glu Gly Arg Glu Leu Leu Leu Cys Ser
485 490 495

Asn Thr Ser Cys Cys Arg Cys Phe Cys Val Glu Cys Leu Glu Val Leu
500 505 510

Val Gly Ala Gly Thr Ala Glu Asp Ala Lys Leu Gln Glu Pro Trp Ser
515 520 525

Cys Tyr Met Cys Leu Pro Gln Arg Cys His Gly Val Leu Arg Arg Arg
530 535 540

Lys Asp Trp Asn Met Arg Leu Gln Asp Phe Phe Thr Thr Asp Pro Asp
545 550 555 560

Leu Glu Glu Phe Glu Pro Pro Lys Leu Tyr Pro Ala Ile Pro Ala Ala
565 570 575

Lys Arg Arg Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala Thr
580 585 590

Gly Tyr Leu Val Leu Lys Glu Leu Gly Ile Lys Val Glu Lys Tyr Ile
595 600 605

Ala Ser Glu Val Cys Ala Glu Ser Ile Ala Val Gly Thr Val Lys His
610 615 620

Glu Gly Gln Ile Lys Tyr Val Asn Asp Val Arg Lys Ile Thr Lys Lys
625 630 635 640

Asn Ile Glu Glu Trp Gly Pro Phe Asp Leu Val Ile Gly Gly Ser Pro
645 650 655

Cys Asn Asp Leu Ser Asn Val Asn Pro Ala Arg Lys Gly Leu Tyr Glu
660 665 670

Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr His Leu Leu Asn Tyr Thr
675 680 685

Arg Pro Lys Glu Gly Asp Asn Arg Pro Phe Phe Trp Met Phe Glu Asn
690 695 700

Val Val Ala Met Lys Val Asn Asp Lys Lys Asp Ile Ser Arg Phe Leu
705 710 715 720

Ala Cys Asn Pro Val Met Ile Asp Ala Ile Lys Val Ser Ala Ala His
725 730 735

Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly Met Asn Arg Pro Val
740 745 750

Met Ala Ser Lys Asn Asp Lys Leu Glu Leu Gln Asp Cys Leu Glu Phe
755 760 765

Ser Arg Thr Ala Lys Leu Lys Lys Val Gln Thr Ile Thr Thr Lys Ser
770 775 780

Asn Ser Ile Arg Gln Gly Lys Asn Gln Leu Phe Pro Val Val Met Asn
785 790 795 800

Gly Lys Asp Asp Val Leu Trp Cys Thr Glu Leu Glu Arg Ile Phe Gly
805 810 815

Phe Pro Ala His Tyr Thr Asp Val Ser Asn Met Gly Arg Gly Ala Arg
820 825 830

Gln Lys Leu Leu Gly Arg Ser Trp Ser Val Pro Val Ile Arg His Leu
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Phe Ala Pro Leu Lys Asp Tyr Phe Ala Cys Glu
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35 40 45

Gly Arg Pro Gly Arg Lys Arg Lys His Pro Pro Val Glu Ser Gly Asp
50 55 60

Thr Pro Lys Asp Pro Ala Val Ile Ser Lys Ser Pro Ser Met Ala Gln
65 70 75 80

Asp Ser Gly Ala Ser Glu Leu Leu Pro Asn Gly Asp Leu Glu Lys Arg
85 90 95

Ser Glu Pro Gln Pro Glu Glu Gly Ser Pro Ala Gly Gly Gln Lys Gly
100 105 110

Gly Ala Pro Ala Glu Gly Glu Gly Ala Ala Glu Thr Leu Pro Glu Ala
115 120 125

Ser Arg Ala Val Glu Asn Gly Cys Cys Thr Pro Lys Glu Gly Arg Gly
130 135 140

Ala Pro Ala Glu Ala Gly Lys Glu Gln Lys Glu Thr Asn Ile Glu Ser
145 150 155 160

Met Lys Met Glu Gly Ser Arg Gly Arg Leu Arg Gly Gly Leu Gly Trp
165 170 175

Glu Ser Ser Leu Arg Gln Arg Pro Met Pro Arg Leu Thr Phe Gln Ala
180 185 190

Gly Asp Pro Tyr Tyr Ile Ser Lys Arg Lys Arg Asp Glu Trp Leu Ala
195 200 205

Arg Trp Lys Arg Glu Ala Glu Lys Lys Ala Lys Val Ile Ala Gly Met
210 215 220

Asn Ala Val Glu Glu Asn Gln Gly Pro Gly Glu Ser Gln Lys Val Glu
225 230 235 240

Glu Ala Ser Pro Pro Ala Val Gln Gln Pro Thr Asp Pro Ala Ser Pro
245 250 255

Thr Val Ala Thr Thr Pro Glu Pro Val Gly Ser Asp Ala Gly Asp Lys
260 265 270

Asn Ala Thr Lys Ala Gly Asp Asp Glu Pro Glu Tyr Glu Asp Gly Arg
275 280 285

Gly Phe Gly Ile Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe Ser
290 295 300

Trp Trp Pro Gly Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser Arg
305 310 315 320

Ala Ala Glu Gly Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys Phe
325 330 335

Ser Val Val Cys Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys Ser
340 345 350

Ala Phe His Gln Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys Ala
355 360 365

Ile Tyr Glu Val Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu Phe
370 375 380

Pro Val Cys His Asp Ser Asp Glu Ser Asp Thr Ala Lys Ala Val Glu
385 390 395 400

Val Gln Asn Lys Pro Met Ile Glu Trp Ala Leu Gly Phe Gln Pro
405 410 415

Ser Gly Pro Lys Gly Leu Glu Pro Pro Glu Glu Lys Asn Pro Tyr
420 425 430

Lys Glu Val Tyr Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala Tyr
435 440 445

Ala Pro Pro Pro Ala Lys Lys Pro Arg Lys Ser Thr Ala Glu Lys
450 455 460

Pro Lys Val Lys Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu Val
465 470 475 480

Tyr Glu Val Arg Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile Ser
485 490 495

Cys Gly Ser Leu Asn Val Thr Leu Glu His Pro Leu Phe Val Gly Gly
500 505 510

Met Cys Gln Asn Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln Tyr
515 520 525

Asp Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly Arg
530 535 540

Glu Val Leu Met Cys Gly Asn Asn Asn Cys Cys Arg Cys Phe Cys Val
545 550 555 560

Glu Cys Val Asp Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala Ile
565 570 575

Lys Glu Asp Pro Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr Tyr
580 585 590

Gly Leu Leu Arg Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met Phe
595 600 605

Phe Ala Asn Asn His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr Pro
610 615 620

Pro Val Pro Ala Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu Phe
625 630 635 640

Asp Gly Ile Ala Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile Gln
645 650 655

Val Asp Arg Tyr Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr Val
660 665 670

Gly Met Val Arg His Gln Gly Lys Ile Met Tyr Val Gly Asp Val Arg
675 680 685

Ser Val Thr Gln Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu Val
690 695 700

Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala Arg
705 710 715 720

Lys Gly Leu Tyr Glu Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr Arg
725 730 735

Leu Leu His Asp Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe Phe
740 745 750

Trp Leu Phe Glu Asn Val Val Ala Met Gly Val Ser Asp Lys Arg Asp
755 760 765

Ile Ser Arg Phe Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys Glu
770 775 780

Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly
785 790 795 800

Met Asn Arg Pro Leu Ala Ser Thr Val Asn Asp Lys Leu Glu Leu Gln
805 810 815

Glu Cys Leu Glu His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg Thr
820 825 830

Ile Thr Thr Arg Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His Phe
835 840 845

Pro Val Phe Met Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu Met
850 855 860

Glu Arg Val Phe Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn Met
865 870 875 880

Ser Arg Leu Ala Arg Gln Arg Leu Leu Gly Arg Ser Trp Ser Val Pro
885 890 895

Val Ile Arg His Leu Phe Ala Pro Leu Lys Glu Tyr Phe Ala Cys Val
900 905 910

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Asp Ser Pro Pro Ile Leu Glu Ala Ile Arg Thr Pro Glu Ile Arg Gly
35 40 45

Arg Arg Ser Ser Ser Arg Leu Ser Lys Arg Glu Val Ser Ser Leu Leu
50 55 60

Ser Tyr Thr Gln Asp Leu Thr Gly Asp Gly Glu Asp Gly Asp
65 70 75 80

Gly Ser Asp Thr Pro Val Met Pro Lys Leu Phe Arg Glu Thr Arg Thr
85 90 95

Arg Ser Glu Ser Pro Ala Val Arg Thr Arg Asn Asn Asn Ser Val Ser
100 105 110

Ser Arg Glu Arg His Arg Pro Ser Pro Arg Ser Thr Arg Gly Arg Gln
115 120 125

Gly Arg Asn His Val Asp Glu Ser Pro Val Glu Phe Pro Ala Thr Arg

130 135 140
Ser Leu Arg Arg Arg Ala Thr Ala Ser Ala Gly Thr Pro Trp Pro Ser
145 150 155 160

Pro Pro Ser Ser Tyr Leu Thr Ile Asp Leu Thr Asp Asp Thr Glu Asp
165 170 175

Thr His Gly Thr Pro Gln Ser Ser Ser Thr Pro Tyr Ala Arg Leu Ala
180 185 190

Gln Asp Ser Gln Gln Gly Gly Met Glu Ser Pro Gln Val Glu Ala Asp
195 200 205

Ser Gly Asp Gly Asp Ser Ser Glu Tyr Gln Asp Gly Lys Glu Phe Gly
210 215 220

Ile Gly Asp Leu Val Trp Gly Lys Ile Lys Gly Phe Ser Trp Trp Pro
225 230 235 240

Ala Met Val Val Ser Trp Lys Ala Thr Ser Lys Arg Gln Ala Met Ser
245 250 255

Gly Met Arg Trp Val Gln Trp Phe Gly Asp Gly Lys Phe Ser Glu Val
260 265 270

Ser Ala Asp Lys Leu Val Ala Leu Gly Leu Phe Ser Gln His Phe Asn
275 280 285

Leu Ala Thr Phe Asn Lys Leu Val Ser Tyr Arg Lys Ala Met Tyr His
290 295 300

Ala Leu Glu Lys Ala Arg Val Arg Ala Gly Lys Thr Phe Pro Ser Ser
305 310 315 320

Pro Gly Asp Ser Leu Glu Asp Gln Leu Lys Pro Met Leu Glu Trp Ala
325 330 335

His Gly Gly Phe Lys Pro Thr Gly Ile Glu Gly Leu Lys Pro Asn Asn

340

345

350

Thr Gln Pro Val Val Asn Lys Ser Lys Val Arg Arg Ala Gly Ser Arg
355 360 365

Lys Leu Glu Ser Arg Lys Tyr Glu Asn Lys Thr Arg Arg Arg Thr Ala
370 375 380

Asp Asp Ser Ala Thr Ser Asp Tyr Cys Pro Ala Pro Lys Arg Leu Lys
385 390 395 400

Thr Asn Cys Tyr Asn Asn Gly Lys Asp Arg Gly Asp Glu Asp Gln Ser
405 410 415

Arg Glu Gln Met Ala Ser Asp Val Ala Asn Asn Lys Ser Ser Leu Glu
420 425 430

Asp Gly Cys Leu Ser Cys Gly Arg Lys Asn Pro Val Ser Phe His Pro
435 440 445

Leu Phe Glu Gly Leu Cys Gln Thr Cys Arg Asp Arg Phe Leu Glu
450 455 460

Leu Phe Tyr Met Tyr Asp Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Val
465 470 475 480

Cys Cys Glu Gly Arg Glu Leu Leu Leu Cys Ser Asn Thr Ser Cys Cys
485 490 495

Arg Cys Phe Cys Val Glu Cys Leu Glu Val Leu Val Gly Thr Gly Thr
500 505 510

Ala Ala Glu Ala Lys Leu Gln Glu Pro Trp Ser Cys Tyr Met Cys Leu
515 520 525

Pro Gln Arg Cys His Gly Val Leu Arg Arg Arg Lys Asp Trp Asn Val
530 535 540

Arg Leu Gln Ala Phe Phe Thr Ser Asp Thr Gly Leu Glu Tyr Glu Ala

545 550 555 560
Pro Lys Leu Tyr Pro Ala Ile Pro Ala Ala Arg Arg Arg Pro Ile Arg
565 570 575

Val Leu Ser Leu Phe Asp Gly Ile Ala Thr Gly Tyr Leu Val Leu Lys
580 585 590

Glu Leu Gly Ile Lys Val Gly Lys Tyr Val Ala Ser Glu Val Cys Glu
595 600 605

Glu Ser Ile Ala Val Gly Thr Val Lys His Glu Gly Asn Ile Lys Tyr
610 615 620

Val Asn Asp Val Arg Asn Ile Thr Lys Lys Asn Ile Glu Glu Trp Gly
625 630 635 640

Pro Phe Asp Leu Val Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Asn
645 650 655

Val Asn Pro Ala Arg Lys Gly Leu Tyr Glu Gly Thr Gly Arg Leu Phe
660 665 670

Phe Glu Phe Tyr His Leu Leu Asn Tyr Ser Arg Pro Lys Glu Gly Asp
675 680 685

Asp Arg Pro Phe Phe Trp Met Phe Glu Asn Val Val Ala Met Lys Val
690 695 700

Gly Asp Lys Arg Asp Ile Ser Arg Phe Leu Glu Cys Asn Pro Val Met
705 710 715 720

Ile Asp Ala Ile Lys Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp
725 730 735

Gly Asn Leu Pro Gly Met Asn Arg Pro Val Ile Ala Ser Lys Asn Asp
740 745 750

Lys Leu Glu Leu Gln Asp Cys Leu Glu Tyr Asn Arg Ile Ala Lys Leu

755

760

765

Lys Lys Val Gln Thr Ile Thr Thr Lys Ser Asn Ser Ile Lys Gln Gly

770

775

780

Lys Asn Gln Leu Phe Pro Val Val Met Asn Gly Lys Glu Asp Val Leu

785

790

795

800

Trp Cys Thr Glu Leu Glu Arg Ile Phe Gly Phe Pro Val His Tyr Thr

805

810

815

Asp Val Ser Asn Met Gly Arg Gly Ala Arg Gln Lys Leu Leu Gly Arg

820

825

830

Ser Trp Ser Val Pro Val Ile Arg His Leu Phe Ala Pro Leu Lys Asp

835

840

845

Tyr Phe Ala Cys Glu

850

<210> 9

<211> 393

<212> DNA

<213> Mus musculus

<400> 9

tttctacagt atttcaggtg cctaccacac aggaaaacctt gaagaaaaacc agtttctaga 60
agccgcgttt acctcttggtt tacagtttat atatatatga tagatatgag atatatata 120
ataaaaggta ctgttaacta ctgtacatcc cgacttcata atggtgcttt caaaacagcg 180
agatgagcaa agacatcagc ttccgcctgg ccctcgtgtg caaatggcgt ttcatgccca 240
tggatggtgt agaggggagc agctggaggg ggttcacaa actgaaggat gacccatatac 300
accccccacc cctgccccat gcctagcttc acctgccaaa aaggggctca gctgaggtgg 360
tcggaccctg gggaaagctga gtgtggaatt tat 393

<210> 10

<211> 424

<212> DNA

<213> Mus musculus

<400> 10
gaagaaaacc agtttctaga agccogctgtt acctcttgc tacagtttat atatatatga 60
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atggtgctt caaaacagcg agatgagcaa agacatcagc ttccgcctgg ccctctgtgc 180
aaagggtttc agcccaggat ggtgagaggg gagcatctgg agggggttt aacaaactga 240
aggatgaccc atatcacccc ccacccctgc cccatgccta gcttcacctg ccaaaaagg 300
gctcagctga ggtggtcgga ccctgggaa gctgagtgtg gaatttatcc agactcgct 360
gcaataacct tagaatatga atctaaaatg actgcctcag aaaaatggct tgagaaaaca 420
ttgt 424

<210> 11
<211> 461
<212> DNA
<213> Mus musculus

<400> 11
tttaaagcaa accacagagg aggaaaacgc cggaggcttg gccttgcaaa agggttggac 60
atcatctcct gagtttcaa tgttaacctt cagtcctatc taaaaagcaa aataggcccc 120
tccccttcgt tcccctccgg tccttaggagg cgaactttt gttttctact cttttcaga 180
ggggttttct gtttggggg gttttgttt cttgctgtga ctgaaacaag agagttattg 240
cagcaaaatc agtaacaaca aaaagttagaa atgccttggc gcggaaagg 300
aattctataa aaacttaaaa tattggttt ttttttttc cttttctata tatctcttg 360
gttgtctcta gcctgatcag ataggagcac aaacaggaag agaatagaga ccctcgagg 420
cagagtctcc tctcccaccc cccgagcagt ctcaacagca c 461

<210> 12
<211> 465
<212> DNA
<213> Mus musculus

<400> 12
tcagaggggt tttctgtttt tttgggtttt tgtttcttgc tgtgactgaa acaagagagt 60
tattgcagca aaatcagtaa caacaaaaag tagaaatgcc ttggagagga aagggagaga 120
ggaaaaattc tataaaaact taaaatattg gttttttttt tttttccctt tctatataatc 180
tctttggttt tctctagcct gatcagatag gagcacaaac aggaagagaa tagagaccct 240
cgaggcaga gtctcccttc ccacccccc agcagtctca acagcaccat tcctggtcat 300
gcaaaacaga acccaactag cagcaggcgc ctgagagaac accacaccag acactttct 360
acagtatttc aggtgcctac cacacagggaa accttgaaga aaaccagttt ctagaagccg 420

ctgttacctc ttgttacag tttatata tatgatagat atgag 465

<210> 13

<211> 393

<212> DNA

<213> Mus musculus

<400> 13

aaaacgccgg aggccttgc cttgcacaag ggttggacat catctcctga gtttcaatg 60
ttaaccttca gtccttatcta aaaagcaaaa taggcccctc cccttcttcc cctccggtcc 120
taggaggcga acttttgtt ttctactctt tttcagaggg gttttctgtt tgtttgggtt 180
tttgggttctt gctgtactg aaacaagaga gttattgcag caaaatcagt aacaacaaaa 240
agtagaaatg cttggagag gaaagggaga gagggaaaat tctataaaaa cttaaaaatat 300
tggttttttt tttttcctt ttctatatacg tctttgggtt gtctctagcc tgatcagata 360
ggagcacaaa caggaagaga atagagaccc tcg 393

<210> 14

<211> 309

<212> DNA

<213> Mus musculus

<400> 14

tgatgattt acgccaaga agtgtctgct gcacacaggg cccgttactt ctagggtaa 60
cttcctggc atgaacaggg ctttggatcc actgtgaatg ataagctgga gctgcaagag 120
tgtctggagc acggcagaat agccaagtgc agcaaagtga ggaccattac caccaggtca 180
aactctataa agcaggcga agaccagcat ttccccgtct tcatgaacga gaaggaggac 240
atcctgtggt gcaactgaaat ggaaagggtc tttggcttcc ccgtccacta cacagacgtc 300
tccaacatg 309

<210> 15

<211> 341

<212> DNA

<213> Mus musculus

<400> 15

tgttaacctt cagtccatc taaaaagcaa aataggcccc tcccccttctt ccctccgg 60
ccttaggagc gaacttttg ttttctactc ttttcagag gggtttctg tttgtttggg 120
ttttgtttc ttgctgtgac tgaaaacaaga gagttattgc agcaaaatca gtaacaacaa 180
aaagttagaaa tgccttggag aggaaaggaa gagagggaaa attctataaa aacttaaaaat 240

atgggtttt tttttttcc ttttctatat atctctttgg ttgtctctag cctgatcaga 300
taggagcaca aacagaaga gaatagagac cctcgaggc a 341

<210> 16
<211> 240
<212> DNA
<213> Mus musculus

<220>
<221> Unsure
<222> (32)..(32)
<223> May be any nucleic acid

<400> 16
acatTTTgta tgTTTTTta tttgctccag gnggggttaa tggcgggtca ctTTCCTca 60
ctctggaata tttctgatcc cacaagggc cttcaacgtg gctgacgaat tcaaaatcag 120
ggacaatgtt ttctcaagcc attttctga ggcagtcat ttagattcat attctaagg 180
tattgcacgc gagtctggat aaattccaca ctcagcttcc ccagggtccg accacctcag 240

<210> 17
<211> 256
<212> DNA
<213> Mus musculus

<220>
<221> Unsure
<222> (75)..(75)
<223> May be any nucleic acid

<400> 17
atcagcttcc gcctggccct ctgtcaaag ggTTcagcc caggatgggg agaggggagc 60
agctggaggg ggttnaaca aactgaaggta tgaccatata caccccccac ccctgccccca 120
tgcctagctt cacctgccaa aaaggggctc agctgaggtg gtcggaccct gggaaagctg 180
agtgtggaat ttatccagac tcgcgtgcaa taaccttaga atatgaatct aaaatgactg 240
cctcagaaaa atggct 256

<210> 18
<211> 435
<212> DNA
<213> Mus musculus

<400> 18

gtggaagccc atgcaatgat ctctctaacg tcaatcctgc ccgcaaagg 60
gcacaggaag gctcttcttc gagtttacc acttgctgaa ttataccgc cccaaggagg 120
gcgacaaccg tccattcttc tggatgttcg agaatgtgt ggccatgaaa gtgaatgaca 180
agaaaagacat ctcaagattc ctggcatgta accoagtgat gatcgatgcc atcaagggt 240
ctgctgctca cagggcccg tacttctggg gtaacctacc cggaatgaac aggcccgtga 300
tggcttcaaa gaatgataag ctcgagctgc aggactgcct ggagttcagt aggacagcaa 360
agttaaagaa agtgcagaca ataaccacca agtcgaactc catcagacag ggcaaaaacc 420
agctttcccc tgtag 435

<210> 19

<211> 522

<212> DNA

<213> Mus musculus

<400> 19

gatgatgtca gcagggatga catcaccacc tttagggctt ttccctggca ggggccccatg 60
tggctagtcc tcacgaagac tggagtagaa tggttggagc tcaggaaggg tgggtggagt 120
ggagtctctt ccaggtgtga gggatacgaa ggaggaagct tagggaaatc cattccccac 180
tccctcttgc caaatgaggg gcccagtc caacagctca ggtccccaga acccccstag 240
tcctcatgag aagcttaggac cagaagcaca tcgttccct tatctgagca gtgtttgggg 300
aactacagtg aaaaccttct ggagatgtta aaagctttt accccacgat agattgtgtt 360
tttaagggggt gctttttta ggggcatcac tggagataag aaagctgcat ttcagaaatg 420
ccatcgtaat gttttttaaa cacctttac ctaattacag gtgctatTTT atagaagcag 480
acaacacttc ttttatgac tctcagactt ctatTTTcat gt 522

<210> 20

<211> 348

<212> DNA

<213> Mus musculus

<400> 20

aaaggaggcc cattagagtc ctgtctctgt ttgatggaaat tgcaacgggg tacttggtgc 60
tcaaggagtt gggtattaaa gtggaaaagt acattgcctc cgaagtctgt gcagagtcca 120
tcgctgtggg aactgttaag catgaaggcc agatcaaata tgtcaatgac gtccggaaaa 180
tcaccaagaa aaatattgaa gagtggggcc cggtcgactt ggtgatttgtt ggaagcccat 240
gcaatgatct ctcttaacgac aatcctgccc gcaaaggTTT atatgagggc acaggaaggc 300
tcttcttcga gttttaccac ttgctgaatt ataccggccc caaggagg 348

<210> 21

<211> 258

<212> DNA

<213> Mus musculus

<400> 21

gttatggtt taagtcttcc tggcaccttc cccttgcttt ggtacaaggg ctgaagtct 60
gttgtcttg tagcatttcc caggatgtat atgtcagcag ggatgacatc atcacctta 120
gggcctttcc ctggcagggg cccatgtggc tagtccac gaagactgga gtataatgtt 180
tggagcttagtgaagggtggg tggagtgtgc ctcttccagg tgtgaggat acgaaggagg 240
aagcttaggg aaatccat 258

<210> 22

<211> 334

<212> DNA

<213> Mus musculus

<400> 22

tgggttaacc tacccgaaat gaacagttaa agaaaagtgc gacaataacc accaagtgc 60
actccatcag acagggcaaa aaccagctt tccctgttagt catgaatggc aaggacgacg 120
ttttgtgttg cactgagctc gaaaggatct tcggcttccc tgctcaactac acggacgtgt 180
ccaaacatggg ccgcggcgcc cgtcagaage tgctggcag gtcctggagt gtacccgtca 240
tcagacacct gtttgcggcc ttgaaggact actttgcctg tgaatagttc tacccaggac 300
tggggagctc tcggtcagag ccagtgc 334

<210> 23

<211> 299

<212> DNA

<213> Mus musculus

<220>

<221> Unsure

<222> (59)..(59)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (173)..(173)

<223> May be any nucleic acid

<400> 23

ctgttttgt ttgtttttt ggtatcttag ccatcacttc tgagtgataa actcaggang 60

gtaaaagaaa gccatcttac tacctacttc aagtttaaa gttcagggt aagagaacat 120
gagcaccatg cgggctact ctaagcagcc aggtctgagc tgtgcacacg ganggagcac 180
cgggctccc ctgcaaggcc aggaggctct gctcccactg agcaggagaa agctgaggt 240
cagtatgtg aggccccaca caggtgagct aaaaaggga caggtgaggt gccttcagg 299

<210> 24

<211> 455

<212> DNA

<213> Mus musculus

<400> 24

gatcgcttcc tagagctctt ctacatgtat gatgaggacg gctatcagtc ctactgcacc 60
gtgtctgtga gggccgtgaa ctgctgctgt gcagtaaacac aagctgctgc agatgcttct 120
gtgtggagtg tctggaggtg ctggggcg caggacagct gaggatgcc a gctgcagga 180
accctggagc tgctatatgt gcctccctca gcgcgtccat ggggtcctcc gacgcaggaa 240
agattgaaac atgcgcctgc aagacttctt cactactgat cctgacctgg aagaatttca 300
ggagccaccc aagttgtacc cagcaattcc tgcagccaaa aggaggccca ttagagtcc 360
gtctctgttt gatgaaatttca acggggta cttggtgctc aaggagttgg gtattaaagt 420
gaaaaagtac attgcctccg aagtctgtgc agagt 455

<210> 25

<211> 368

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (307)..(307)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (335)..(335)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (353)..(353)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (360)..(360)

<223> May be any nucleic acid

<400> 25
acgtttgtatgttttta tttgtccag gtggggttt gactgtcact ttcccacact 60
ctggattagt tctgatccca ccacaaggag ccctcgaaatt ggctaaagtg agaaaactggg 120
cctgaagact ccgtaccctc tgccatcttgc cgagggagt ctcccttttag aaaacaatca 180
aagggttatt gcatgagtctt gatgaatcc cacttcagc ttgtccacgg gcccgaccac 240
ctcatctagc ccccttttg gcaagggaga acctggctcc caagttctcc tccttcaactt 300
tcgttancaa accaaggggg aagaagccca ccgtngagaa cgcgccatct tgnaaagctn 360
ggtcttcc 368

<210> 26

<211> 399

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (87)..(87)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (314)..(314)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (318)..(318)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (370)..(370)

<223> May be any nucleic acid

<400> 26

gaacatgagg atggagagaa gtatcagcac ccagaagaga aaaaggaatt taaaacaaaa 60
accacagagg cgaaaatacc ggaggcntt gcttgcgaaa agggttggac atcatctcct 120
gatTTTCAA TGTATTCTT CAGTCCTATT TAAAAACAAA ACCAAGCTCC CTTCCTTCC 180
TCCCCCTTCC CTTTTTTC GGTCAAGACCT TTTATTTCT ACTCTTTCA GAGGGTTTT 240
CTGTTTGTAA GGGTTTGTG TCTTGCTGTG ACTGAAACAA GAAGGTTATT GCAGCAAAAA 300
TCAGGTAACA AANATANGT AACAATACCT TGCAAGAGGA AGGTGGGAGG AGAGGAAAAA 360
AGGGAAATTN CTATAGAAAT CTATATATTG GGTTGGTT 399

<210> 27

<211> 318

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (205)..(205)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (275)..(275)

<223> May be any nucleic acid

<400> 27

gtacgaggtg cggcagaagt gccggaacat tgaggacatc tgcatactcct gtgggagcct 60
caatgttacc ctggaacacc ccctttcggt tggaggaatg tgccaaaact gcaagaactg 120
ctttctggag tgtgcgtacc agtacgacga cgacggctac cagtcctact gcaccatctg 180
ctgtgggggc cgtgaggtgc tcatntgcgg aaacaacaac tgctgcaggt gctttgcgt 240
ggagtgtgtg gacctcttgg tggggccggg ggctncccag gcagcagtta aggaagatca 300
tgtacgtcgg ggacgtcc 318

<210> 28

<211> 259

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (227)..(227)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (234)..(234)

<223> May be any nucleic acid

<400> 28

gagccgagca gctgaaggca cccgctgggt catgtggttc ggagacggca aattctcagt 60
ggtgtgtgtt gagaagctga tgccgctgag ctcgtttgc agtgcgttcc accaggccac 120
gtacaacaag cagccatgtt accgcaaagc catctacgag gtcctgcagg tggccagcag 180
ccgcgcgggg aagctgttcc cggtgtgcca cgacagcgat gagagtnaca ctgncaaggc 240
cgtgggaggt gcagaacaa 259

<210> 29

<211> 483

<212> DNA

<213> Homo sapiens

<400> 29

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acactctgga ttagttctga tcccaccaca aggagccctc gaattggcta aagtgagaaa 120
ctgggcctga agactccgta ccctctgcca tcttgcggag ggagtctctt tttagaaaaac 180
aatcaaaggg ttattgcatg agtctggatg aatcccactc tcagctgtcc acggggccga 240
ccacacctatc taggccccctt tttggcaagg agaacccggg tcccaagttc tcctccttca 300
cttcgttaca aaccaggggg aaaaagccca cgtaaaaacg cgccatctgc aaaatggttc 360
cccttcttca tccctgggaa aacctttgcg ccaaggcaac gtggaaaactg atggtttac 420
tcaactcgct gtttgaagc gccattatga aatcggggtt gtacgtaggt aaagtcccgt 480
gcc 483

483

<210> 30

<211> 337

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (41) .. (41)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (45) .. (45)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (176) .. (176)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (190) .. (190)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (207) .. (207)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (265)..(265)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (290)..(290)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (317)..(317)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (322)..(322)
<223> May be any nucleic acid

<400> 30
gggcattcag gtggaccgct acattgcctc ggaggtgtgt naggncatcca tcacggtggg 60
catggtcgg caccaggaga agatcatgtc cgtagggac gtccgcacgc tcacacagaa 120
gcataatccag gagtgccgc cattcgatct ggtgattggg ggcagtccct gcaatnacct 180
ctccatcgtn aaccctgctc gcaaggncct ctacgagggc actggccggc tcttctttaa 240
gttctaccgc ctccatcgatg atgcncggcc caaggaggaa agatgatcgn cccttcttct 300
ggctctttaa gaatgtngtg gnccatggc gtttagt 337

<210> 31
<211> 271
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (234)..(234)
<223> May be any nucleic acid

<400> 31
cttgtttaca gtttatatat atatgataga tatgagatat atatataaaa ggtactgtta 60
actactgtac aacccgactt cataatggtg ctttcaaaca gcgagatgag taaaaacatc 120
agcttccacg ttgccttctg cgcaaagggt ttcaccaagg atggagaaaag ggagacagct 180
tgcagatggc gcgttctcac ggtgggctct tcccttggt ttgttaacgaa gtgnaggagg 240
agaacttggg agccaggttc tccctgccaa a 271

<210> 32
<211> 430

<212> DNA

<213> Homo sapiens

<400> 32

acgttttgta tgtttttta tttgtccag gtggggttt gactgtact ttccccacact 60
ctggattagt tctgatccc ccacaaggag ccctcgaatt ggctaaagtg agaaaactggg 120
cctgaagact ccgtaccctc tgccatcttgc cgagggagt ctcctttaga aaacaatcaa 180
agggttatttgc catgagtctg gatgaatccc actctcagct gtccacgggc ccgaccacct 240
catctagccc ccttttggc agggagaacc tggctcccaa gttctcctcc ttcacttcgt 300
tacaaacccaa gggaaagagc ccaccgtgag aacgcgcctt ctgcaagctg tctcccttgc 360
tccatccttgc gtgaaacccc tttgcgcaga aggcaacgtg gaagctgatg tttttactca 420
tctcgctgtt 430

430

<210> 33

<211> 483

<212> DNA

<213> Homo sapiens

<400> 33

ttttttttt ttgtatgttt ttttatttgc tccaggtggg gttttgactg tcactttccc 60
acactctgga ttagttctga tcccaccaca aggagccctc gaattggcta aagtgagaaa 120
ctgggcctga agactccgta ccctctgcca tcttgccgag ggagtctctt tttagaaaaac 180
aatcaaaggg ttattgcatg agtctggatg aatcccactc tcagctgtcc acggggccga 240
ccacacctatc taggccccctt tttggcaagg agaaccgggg tcccaagttc tcctccttca 300
cttcgttaca aaccaggggg aaaaagccca cgtaaaaacg cgccatctgc aaaatggttc 360
cccttcttca tccctgggg aacctttgcg ccaaggcaac gtggaaactg atggttttac 420
tcaactcgct gtttgaagc gccatttatga aatcggggtt gtacgttaggt aaagtcccg 480
gcc 483

483

<210> 34

<211> 411

<212> DNA

<213> Homo sapiens

<400> 34

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gaaactgggc ctgaagactc cgtaccctct gccattttcgc cgagggagtc tccttttaga 180
aaacaatcaa agggttatttg catgagtctg gatgaatccc actctcagct gtccacgggc 240

ccgaccacct catctagccc cctttggca gggagaacct ggctcccaag ttctcctcct 300
tcacttcgtt acaaaccagg gggaaagagcc caccgtgaga acgcgccatc tgcaagctgt 360
ctcccttctt ccattttgg taaaaccctt tgccgagaag gcaacgtgga a 411

<210> 35

<211> 530

<212> DNA

<213> Homo sapiens

<400> 35

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ttcgctccgc tgaaggcgta ttttgcgtgt gtctaaggga catggggca aactgaggta 120
gacacacaaa gttaaacaca caaacacccc acacacaaca taataacaaca ccaagaacat 180
gaggatggag agaagtatca gccacccaga agagaacaag gaatttaaaa ccaaaaaccac 240
agaggcggaa ataccggagg actttgcctt gcgaccagg ttggacatca tctcctgatt 300
tttcaatgtt attcttcagt cctatttaaa aacaaaacca agtccccctc ctttcctgcg 360
gctcccttt ttttcggc agaccttttta ttttctactc ttttcagagg ggtttctgt 420
ttgtttgggt tttgtttctt gctgtgactg aaacaagaag gttattgcag caaaaatcag 480
taacaaaaaa tagtaacaat accttgacaga ggaaaggtgg gagagaggaa 530

<210> 36

<211> 535

<212> DNA

<213> Homo sapiens

<400> 36

tttacgtttt gtatgtttt ttatggctc caggtgggt tttgactgtc actttccac 60
actctggatt agttctgatc ccaccacaag gagccctcga attggctaaa gtgagaaact 120
ggccctgaag actccgtacc ctctgccatc ttgccgaggg agtctccctt tagaaaacaa 180
tcaaagggtt attgcatgag tctggatgaa tcccactctc agtgcac 240
acccatcta gccccctttt tggcaggag aacctggctc ccaagttctc ctccttcact 300
tcgttacaaa ccacggggaa gagccacccg tgagaacgcg ccatctgcaa gctgtctccc 360
tttctccatc cttgggtaaa cccttgcgc agaaggcaac gtggaaagctg atgttttac 420
tcatctcgct gtttggaaagc accattatga agtcgggttg tacagttagtt aacagtacct 480
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<210> 37

<211> 428

<212> DNA

<213> Homo sapiens

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<222> (15)..(15)

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<222> (424)..(424)

<223> May be any nucleic acid

<400> 37

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cctgaagact ccgtacccttc tgccatcttgc ccgagggagt ctcccttttag aaaacaatca 180
aagggttatt gcatgagtctt gatatcc cactctcagc tgtccacggg cccgaccacc 240
tcatcttagcc cccttttgg cagggagaac ctgggctccc aagttctccct ctttcacttc 300
gttacaaacc aaggggaagg agcccaccgt gagaacggcg ccatcttgca agctgtctcc 360
ctttctccat ctttgggttga aacccttttgc cgccagaagg caacgtggga agctngatgt 420
tttntaac 428

<210> 38

<211> 419

<212> DNA

<213> Homo sapiens

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atgaacagggc cgttggatcc actgtgaatg ataagctgga gctgcaggag tgtctggagc 180
atggcaggat agccaagttc agcaaagtga ggaccattac tacgaggatca aactccataaa 240
agcaggggcaa agaccagcat tttcctgtct tcatgaatga gaaagaggac atcttatgg 300
gcactnaaat tggaaagggtt atttngggtt tcccagtcca ntatactgac gtctccaaca 360
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<223> May be any nucleic acid

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<221> Unsure

<222> (426)..(426)

<223> May be any nucleic acid

<400> 39

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caactcaggcg gcacanttcc ctcccagcca ctgagctgtc gtgccagcac cattcctgg 120
cacgc当地
cagc当地
tacagtattt caggtgccta ccacacagga aaccttgaag aaantcagtt tcttaggaaggc 240
cgctgttacc tcttgtttac agtttatata tatatgatag atatgagatn tatatataaaa 300
aggtaactgtt aactactgtt caacccgact tcataatggg tgctttcaaa cagggcaggt 360
gngtaaaaac atcagnttcc acgttngcct tttgcgc当地
aaggngaca gctttt 420
437

<210> 40

<211> 385

<212> DNA

<213> Homo sapiens

<220>

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<222> (340)..(340)

<223> May be any nucleic acid

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<221> Unsure

<222> (365)..(365)
<223> May be any nucleic acid

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<223> May be any nucleic acid

<400> 40

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gcctgaagac tccgtaccct ctgccatctt gccgaggggag ttcctttta gaaaacaatc 180
aaagggttat tgcatacgatc tggatgaatc ccactctcag ctgtccacgg gccccgaccac 240
ctcatcttagc ccccttttg gcagggagaa cctgggctcc caagttctcc tccttcaatt 300
cgttacaaac caagggaaag agcccaccgt gagaacgcgn catctgcaag ctgtctccct 360
tttncatcc ttggtrngaaa ccctt 385

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<211> 294
<212> DNA
<213> Homo sapiens

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<222> (73)..(73)
<223> May be any nucleic acid

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<223> May be any nucleic acid

<400> 41

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ttgtctctag cctgatcaga taggagcaca agcaggggac ggaaagagag agacactcag 180
gccccacatt tgccctccca gccactgagc tgtcgtgcca gcaccattcc tgggtcacgc 240
aaaacagaac ccagttagca gcagggnaga cgagaacacc acacaagaca tttt 294

<210> 42
<211> 610
<212> DNA
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<223> May be any nucleic acid

<220>
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<223> May be any nucleic acid

<400> 42
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gcctgaagac tccgtaccct ctgccatctt gccgagggag ttcctttta gaaaacaatc 180
aaagggttat tgcgtgatgc tggatgaatc ccactctcag ctgtccacgg gcccgaccac 240
ctcatcttagc ccccttttg gcagggagaa cctggctccc aagttctctt ctttcacttc 300
gttacaaacc aaggggaaga gcccaccgtg agaacgcgcc atctgcaagc tgtctccctt 360
tctccatcct ttggtgaaaa ccctttgcg cagaaggcaa cgtggaagct gatgtttta 420
ctcatctcgc tggatggaaag caccattatg aagtcgggtt gtacagtagt taacagtacc 480
tttatatat atatctcata tctatcatat atatataaac tggtaaacaa gaggtAACAG 540
cggttttcta gaaactgatt ttcttcaagg tttccngtgt ggtggcacn tgaaatactg 600
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<210> 43
<211> 283
<212> DNA
<213> Homo sapiens

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<223> May be any nucleic acid

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<223> May be any nucleic acid

<400> 43
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tgcctcgcca agcggctcat gttggagacg tcagtagtggact ggactggaa accaaatacc 180
cttccatccat cagtgcacca taagatgtcc tcttcctcat tcataagac aggaaaaatg 240
ctggtcatttgcgccttta tggagtttg anctcgtaag taa 283

<210> 44

<211> 383

<212> DNA

<213> Homo sapiens

<400> 44

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tgattgggggg cagtccctgc aatgacacctt ccatcgtaaa ccctgctcgc aaggggctct 120
acgagggcac tggccggctc ttctttgagt tctaccgcct cctgcgtatgat ggcggccca 180
aggagggaga tgatcgcccc ttctctggct ctggagaat ttggtgccca tggcgtagt 240
acacagagag gacacatctc gcgatttctc gagtccaacc ctgtatatga ttgatgccaa 300
agaagtctca tctgcacaga ggcccctcta cttctgggt cacctccccg tattaacagg 360
ccgtaggatc cactgttatt ata 383

<210> 45

<211> 447

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (445) .. (445)

<223> May be any nucleic acid

<400> 45

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cctgaagact ccgtaccctc tgccatctg ccgagggagt ctccttttag aaaacaatca 180
aagggttatt gcatgagtct ggatgaatcc cactctcagc tgcacggg cccgaccacc 240
tcatactcaagc ccccttttg gcagggagaa cctggctccc aagttctcct ctttcacttc 300
gttacaaacc aaggggaaaga gcccaccgtg agaacgcgcc atctgcaagc tgcacccctt 360
tctccatcct tggtaaaacc ttgcgcaga aggcaacgtg gaaagctgaa gtttttact 420
catctcgctg tttgaaaagc accanta 447

<210> 46

<211> 100

<212> DNA

<213> Homo sapiens

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<222> (96)..(96)

<223> May be any nucleic acid

<400> 46

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aaacaaaaaac cacagaggcg gaaataccgg tgactnttct 100

<210> 47

<211> 150

<212> DNA

<213> Homo sapiens

<400> 47

tactccttca gcgggttagga ggtggcggat gactggcacg ctccatgacc ggcccagcag 60
tctctgcctc gccaaagcgct catgttggag aggtcagttat agtggactgg gaaaccaaat 120
accctttcca tttcagtgca ccataagatg 150

<210> 48

<211> 237

<212> DNA

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<400> 48

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tttacccacc tgtcccaagct gagaagagga agcccatccg ggtgctgtct ctctttgatg 180

gaatcgctac aggtgagggg tgcaggccca agaggtgctg gcctcgtgcg aattcct 237

<210> 49

<211> 442

<212> DNA

<213> Homo sapiens

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<223> May be any nucleic acid

<400> 49

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ccagcaccat tcctggncac gcaaaacaga acccagttag cagcagggag acgagaacac 180
cacacaagac attttctac agtattttag gtgcctacca cacagggaaa cttgaagaaa 240
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tatgaggata tatatataaa agggtactgt ttaactactg taccaacccg actttataaa 360
tgggtgcttt tcaaacagcc gaggatgnng taaaancat cagcttccac gttgccttct 420
gcggcaangg gtttaccag gg 442

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<211> 395

<212> DNA

<213> Homo sapiens

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<400> 50

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gcctgaagac tccgtaccct ctgccatctt gccgagggag ttcctttta gaaaacaatc 180
aaagggttat tgcatgagtc tggatgaatc ccactctcag ctgtccacgg gccccaccac 240
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ttcgtaaca aaccaagggg aagagccac cgtgaggaac ggngccatct ggcaaggttg 360
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<210> 51

<211> 835

<212> DNA

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<223> May be any nucleic acid

<400> 51

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tacctcagtt tgccccatn tccctnacac acacgcaaaa tactccttca gcggagcga 180
gaggtggcgg atgactggna cgctccatga ccggcccgagc agtctctgcc tcgccaagcg 240
gatcatgttg gagacgtcag tatagtggac tggaaacca aatacccttt ccatttcagn 300
gcaccataag atgtcctctt tctcattcat gaagacaggg aaaatgctgg tctttggcct 360
gctcnatgga gtttgactcc gtagtaangg ccctcanttt ggntgacttg ggctatcctg 420
ncatgctcca gacacctccg nagggtcaca acagaagcat ntccagggg gtggnggcca 480
ttccgacctt tggnggattg ggggggaagc cccnaaaaat aacccttca aacggnnnaaa 540
ccctngttcn gaangggccc ctttncgang ggaaactgggn ccgnttnttt cttnngggnt 600
tcctcccccc ccccccnnaaa ataatggng gccccaaagna gggaaattac cccccccncn 660
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ttttggnccc ncccnnnnt tnnnncccaa aaannntaat taaaaaggcc cttttctggg 780
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gcctgaagac tccgtaccct ctgccatctt gccgagggag ttcctttta gaaaacaatc 180
aaagggttat tgcatacgatc tggatgaatc ccactctcag ctgtccacgg gcccgaccac 240
ctcatcttagc ccccttttg gcagggagaa cctggctccc aagttctctt ctttcacttc 300
gttacaaacc aaggagaaga gcccaccatg agaacgcgcc atctgcaagc tgtctccctt 360
tctncatctt tggtaaaacc tttgcgcaga aggcaacgtg gaagctgatg ttttntcat 420
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tagatatgag atatatatataaaaaggtact gttaactact gtacaacccg acttcataat 180

ggtgcttca aacagcgaga tgagtaaaaa catcagcttc cacgttgcct tctgcgcaaa 240
gggttcacc aaggatggag aaaggagac agcttcaga tggcgcttc tcattggtggg 300
ctctccct tggttgtaa cgaagtntag gaggagaact tgggagccag gttctccctg 360
ccaaaaaggg ggcttagatga ggtggtcggg cccgtggaca gctgagagtg ggattcatcc 420
agactcatgc aataaccctt tgattgttc taaaaggaga ctccctcggc aagatggcag 480
agggtacgga gtcttcaggc ccagttntca cttagccaa t 521

<210> 54

<211> 440

<212> DNA

<213> Homo sapiens

<400> 54

ctctcttga tggaatcgct acaggctcc tggtgctgaa ggacttgggc attcaggtgg 60
accgctacat tgcctcggag gtgtgtgagg actccatcac ggtgggcacg gtgcggcacc 120
aggggaagat catgtacgtc ggggacgtcc gcagcgtcac acagaagcat atccaggagt 180
ggggccatt cgatctggtg attgggggca gtccctgcaa tgacctctcc atcgtcaacc 240
ctgctcgcaa gggcctctac gagggcactg gccggctctt cttaggttc taccgcctcc 300
tgcgtatgc gggcccaag gagggagatg atcgccccctt cttctggctc tttgagaatg 360
tggtgccat gggcgtag tgacaagagg gacatctcgc gatttctcga gtccaaccct 420
gtgtatgattt atgccaaga 440

<210> 55

<211> 273

<212> DNA

<213> Homo sapiens

<400> 55

acgtttgtatgttttta ttgtccag gtgggtttt gactgtact ttcccacact 60
ctggattagt tctgatccca ccacaaggag ccctcgaatt ggctaaagtg agaaactggg 120
cctgaagact ccgtaccctc tgccatcttgc cgagggagt ctccctttag aaaacaatca 180
aagggttatt gcatgagtct gatgaatcc cactctcgc tgtccacggg cccgaccacc 240
tcatctagcc cccttttgg cagggagaac ctg 273

<210> 56

<211> 190

<212> DNA

<213> Homo sapiens

<220>
<221> Unsure
<222> (39)..(39)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (83)..(83)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (181)..(181)
<223> May be any nucleic acid

<400> 56
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agaagagaaa aaggaattta aancaaaaac cacagaggcg gaaataccgg agggctttgc 120
cttgcgaaaa gggttggaca tcatctccctg attttcaat gttattcttc agtcctat 180
naaaacaaag 190

<210> 57
<211> 445
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (167)..(167)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (353)..(353)
<223> May be any nucleic acid

<400> 57
tttagacaaat actgattttta attaaacata aggtaaactc taggcattccg tcatctttca 60
gcctaaaaat tagcaaaaac tggtaaaaca aggacacagtt ttttccccat atttgttacg 120
tcgtggctcc agttacaaaa aaattttaat gaaaacgtta aacatanaaa tagaagttt 180
agattttaaa aagtgtataa aaagccccac aaaacttgc aacgggttgc ctttattcta 240
caaaatagca ccagtaagaa gagtaaaagg tgtaaaaac catttatgac agcatttctg 300
aaatgcagct tgtctgaatt cccggttctc cctaaaaacg acttctttat ggnattaaaa 360
aagggtttaa aaaaatctcc aaaggggagc accgagctt gcaggtttc cctgtcatct 420
ctcagatgtg ggggaagctc gtggc 445

<210> 58
<211> 287
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (38)..(38)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (171)..(171)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (204)..(204)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (274)..(274)
<223> May be any nucleic acid

<400> 58
ttccccacat ctgagagatg acaggaaaaa ctgcaaanc tggctccc tttggagatt 60
tttaatcct ttttattcc ataagaagtc gtttttaggg agaacggaa ttca gacaag 120
ctgcattca gaaatgctgt cataatggtt tttaacacct ttactcctc ntta ctttggtg 180
ctattttgt agaataaggg aacnacgtt acaagttt gttttttttt tttatacac 240
ctttttaaa atctccaact tcctaattt taanggtta accgttt 287

<210> 59
<211> 535
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (452)..(452)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (526)..(526)
<223> May be any nucleic acid

<400> 59

tagacaaata ctgatttaa taaaacataa ggtaaactct aggcatccgt catctttcag 60
cctaaaaatt agcaaaaact gttgaaacaa ggcacagttt tttccccata tttgttacgt 120
cgtggctcca gttacaaaaa aattttaatg aaaacgttaa acataaaaat agaagttga 180
gattttaaaa agtgtataaa aagccccaca aaacttgtca acgttgttcc ttattctaca 240
aaataggcacc agtaagaaga gtaaaagggtg ttaaaaacca ttatgacagc atttctgaaa 300
tgcagcttgt ctgaattccc gttctcccta aaaacgactt cttatggaat aaaaaaggat 360
taaaaaatct ccaaagggag caccgagctt tgcagtttc cctgtccgtc tctcagatgt 420
gggaaaggta tgagaaatgt atgtctgtcc cngactgctg tcactgcctc tgagtttagta 480
aaaggtgaga atgaggtag cagcttccca tctgggcct gtgccngtgg aggg 535

<210> 60

<211> 449

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (7)..(7)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (200)..(200)

<223> May be any nucleic acid

<400> 60

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tctgaagtgt gtgaggagtc cattgctgtt ggaaccgtga agcacgaggg gaatatcaaa 120
tacgtgaacg acgtgaggaa catcacaaag aaaaatattg aagaatgggg cccatttgac 180
ttggtgattg gcggaaccan tgcaacgatc tctcaaattgt gaatccagcc aggaaaggcc 240
tgtatgaggg tacaggccgg ctcttcttcg aattttacca cctgctgaat tactcacgcc 300
ccaaggaggg tcatgaccgg ccgttcttct ggatgttga gaatgttga gccatgaagg 360
ttggcgacaa gagggacatc tcacggttcc tggagtgtaa tccagtgtatg attgatgcc 420
tccaaagttt ctgctgctca cagggcccg 449

<210> 61

<211> 522

<212> DNA

<213> Homo sapiens

<220>
<221> Unsure
<222> (146)..(146)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (281)..(281)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (304)..(304)
<223> May be any nucleic acid

<400> 61
aagagggaca tctcacgggtt cctggagtgt aatccagtga tgattgatgc catcaaagtt 60
tctgctgctc acagggccccg atacttctgg ggcaacctac ccgggatgaa caggcccgtg 120
atagcatcaa agaatgataa actcgngctg caggactgct tggaaatacaa taggatagcc 180
aagttaaaga aagtacagac aataaccacc aagtctgaact cgatcaaaca ggggaaaaac 240
caactttcc ctgttgtcat gaatggcaaa gaagatgttt ngtggtgac tgagctcgaa 300
agntcttg gctttcctgt gcactacaca gacgtgtcca acatgggccccg tggtgccgc 360
cagaagctgc tgggaaggtc ctggagcgtg cctgtcatcc gacacctt cgcctctg 420
aaggactact ttgcatgtga atagttccag ccagggccca agcccactgg ggtgtgtggc 480
agagcaggac ccaggaggtg tgattctgaa ggcattccca gg 522

<210> 62
<211> 573
<212> DNA
<213> Homo sapiens

<400> 62
ctaagatcca ttttctaaac tccaatttag cattctctgt atctgggtgg tttttacttt 60
tttacttaat cttgcttgat caggaactct ggtgtcttct tggcccccca cgtgatctcg 120
ttcatggtca cttttttgtt tatctcattt tctctgaggc tggccttcc tggtaacgtc 180
ttggcatttg tgggaagcac aaaatgttct tgtccctcca actctgttt tcgctccctg 240
ccctgccatt cctctcccgcc gcctgccctc tccctccat cttcccaagg tactttctc 300
tcccagccct gccactcttc tgccgcacct ggcgtctccc ctccatctt cccaggtact 360
tttgagcctt gactccccag gtcccttcat tctgtgtca ctccatgtatg tcattttgtt 420
ctccagttaa agaaagtaca gacaataacc accaagtctga actcgatcaa acagggaaa 480
aaccaacttt tccctgttgtt catgaatggc aaagaagatg ttttgggtg cactgagctc 540
gaaaggatct ttggcttcc tgtgcactac aca 573

<210> 63
<211> 559
<212> DNA
<213> Homo sapiens

<400> 63
agacaaatac tgatttaat taaacataag gtaaactcta ggcatccgtc atcttcagc 60
ctaaaaatta gcaaaaactg ttgaaacaag gcacagtttt ttccccatat ttgttacgtc 120
gtggctccag ttacaaaaaa attttaatga aaacgttaaa cataaaaaata gaagtttgag 180
attttaaaaa gtgtataaaa agccccacaa aacttgtcaa cgttgttcct tattctacaa 240
aatagcacca gtaagaagag taaaagggtgt taaaaaccat tatgacagca tttctgaaat 300
gcagcttgc tgaattcccg ttctccctaa aaacgacttc ttatggaata aaaaaggatt 360
aaaaaatctc caaaggggagc accgagcttt gcagtttcc ctgtcatcta tcagatgtgg 420
ggaaggtatg agaaatgtat gtctgtccct gactgctgtc actgcctctg agtttagtaa 480
aaagatgaga aatgagggtta gcagacttct catctgggga cctgtgcctg tggagggtag 540
gtctcctgga gagggaatg 559

<210> 64
<211> 391
<212> DNA
<213> Homo sapiens

<400> 64
tttttttta gacaaatact gatTTtaatt aaacataagg taaactctag gcatccgtca 60
tctttcagcc taaaaattag caaaaactgt tgaacaagg cacagttttt tccccatatt 120
tgttacgtcg tggctccagt tacaaaaaaa attttaatga aaacgttaaa cataaaaaata 180
gaagtttgag attttaaaaa gtgtataaaa agccccacaa aacttgtcaa cgttgttcct 240
tattctacaa aatagcacca gtaagaagag taaaagggtgt taaaaaccat tatgacagca 300
tttctgaaat gcagcttgc tgaattcccg ttctccctaa aaacgacttc ttatggaata 360
aaaaaggatt aaaaatctc caaaggggagc a 391

<210> 65
<211> 517
<212> DNA
<213> Homo sapiens

<400> 65
acaaaatactg attttatTT aacataaggt aaactctagg caggggcattc tttcagccta 60

aaaattagca aaaactgttg aaacaaggca cagtttttc cccatatttgc tacgtcg 120
gctccagtttta cgaaaaattttaatgaaaa cgttaaacat aaaaatagaa gttttagattt 180
ttaaaaagtgtataaaaagc cccacaaaaac ttgtcaacgt tgcccttat tctacaaaat 240
agcaccagta agaagagtaa aagggtttaa aaaccattat gacagcattt ctgaaatgca 300
gcttgtctga attccgttc tccctaaaaa cgacttctta tggaaaaaaa aaggattaaa 360
aaatctccaa agggagcacc gagcttgca gtttccctg tcacatctca gatgtggg 420
aggatgaga aatgtatgtc tgtccctgac tgctgtcact gcctctgagt ttagaaaaa 480
gatgagaaat gagggtagca gacttctcat ctgggg 517

<210> 66

<211> 442

<212> DNA

<213> Homo sapiens

<400> 66

gacaaataact gatttaattt aaacataagg taaactctag gcatccgtca tcttcagcc 60
taaaaaattttt caaaaactgt tgaaacaagg cacagttttt tccccatattt tggtacgtcg 120
tggctccagt tacaaaaaaa attttaatgtt aacgtttaaa cataaaaaata gaagtttgag 180
attttaaaaaa gtgtataaaaaa agccccacaa aacttgcacaa cgttgttccctt tattctacaa 240
aatagcacca gtaagaagag taaaagggtgt taaaaccat tatgacagca tttctgaaat 300
gcagcttgcc tgaattcccg ttctccctaa aaacgacttc ttatggaata aaaaaggattt 360
aaaaaatctc caaaggagc accgagctt gcagtttcc ctgtcatctc gcagatgtgg 420
ggaaggatgtt agaaatgtat gt 442

<210> 67

<211> 396

<212> DNA

<213> Homo sapiens

<400> 67

gcagtcaggg acagacatac atttctcata cttccccac atctgagaga tgacagggaa 60
aactgcacaaag ctcggtgctc ccttggaga tttttatc cttttttt ccataagaag 120
tcgttttag ggagaacggg aattcagaca agctgcattt cagaaatgct gtcataatgg 180
tttttaacac ctttactct tcttactgtt gctatgtt agaataagga acaacgttga 240
caagtttgc ggggctttt atacactttt taaaatctca aacttctatt tttatgttta 300
acgttttcat taaaatttt ttgtactgg agccacgacg taacaaatat gggaaaaaaa 360
ctgtgccttgc tttcaacagt ttttgcataat ttttag 396

<210> 68

<211> 287

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (7)..(7)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (169)..(169)

<223> May be any nucleic acid

<400> 68

agacaantac tgatttaat taaacataag gtaaactcta ggcatccgtc atcttcagc 60
ctaaaaatta gcaaaaactg ttgaaacaag gcacagttt tccccatat ttgttacgtc 120
gtggctccag ttacaaaaaa aattttaatg aaaacgttaa acataaaaant agaagtttga 180
gatttaaaaa agtgtataaa aagccccaca aaacttgtca acgttgttcc ttattctaca 240
aaatagcacc agtaagaaga gtaaaagggtg ttaaaaacca ttatgac 287

<210> 69

<211> 356

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (193)..(193)

<223> May be any nucleic acid

<400> 69

attgaagaat gggcccatt tgacttggtg attggcgaa ccgatgcaac gatctctcaa 60
atgtaatcc agccaggaaa gcctgtatg agggtacagg ccggcttcc ttcaatttt 120
accacctgct gaattactca cgccccaagg agggtgatga ccggccgttc ttctggatgt 180
ttgagaatgt tgnagccatg aaggttggcg acaagaggga catctcacgg ttccctggagt 240
gtaatccagt gatgattgat gccatcaaag tttctgctgc tcacagggcc cgataactct 300
ggggcaacct acccgggatg aacaggatct ttggctttcc tgtgcactac acagac 356

<210> 70

<211> 408

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (408)..(408)

<223> May be any nucleic acid

<400> 70

ttagacaaa tactgattt aattaaacat aaggtaaact ctaggcattcc gtcatcttc 60
agcctaaaaa ttagcaaaaaa ctgttgaaac aaggcacagt ttttccccca tatttggta 120
gtcggtggc cagttacaaa aaaaattttt atgaaaacgt taaacataaa aatagaagtt 180
ttagattttt aaaaagtgtat aaaaagcccc acaaaaacttg tcaacgttgt tccttattct 240
acaaaaatagc accagtaaga agagtaaaag gtgttaaaaa ccattatgac agcatttctg 300
aatatcgagct tgtctgaatt cccgttctcc ctaaaaaacga cttcttatgg aataaaaaaag 360
gattaaaaaaaaa tctccaaagg gagcaccgag ctttgcagtt ttccctgn 408

<210> 71

<211> 439

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (50)..(50)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (85)..(85)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (405)..(405)

<223> May be any nucleic acid

<400> 71

gcatgttagct acaggacatt tttaagggcc caggatcggt ttttcccagn tgcaagcaga 60
agagaaaatg ttgtatatgt ctttnaccgg gcacattccc cttgcctaaa tacaagggt 120
ggagtctgca cgggacctat tagagtattt tccacaatga tgatgatttc agcagggatg 180
acgtcatcat cacattcagg gctattttt cccccacaaa cccaaggcga gggccactc 240
ttagctaaat ccctccccgt gactgcaata gaaccctctg gggagctcag gaaagggggt 300
gtgctgagtt ctataatata agctgccata tattttgtag acaagtatgg ctcctccat 360

atctccctct tccctaggag aggagtgtga aagcaaggga gcttngataa gacaccccct 420
caaaccatt ccctctcca 439

<210> 72
<211> 491
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (26)..(27)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (33)..(33)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (188)..(188)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (301)..(301)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (339)..(339)
<223> May be any nucleic acid

<220>
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<222> (360)..(360)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (379)..(379)
<223> May be any nucleic acid

<400> 72
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aaactgttga aacaaggcac agtttttcc ccataattgt tacgtcgtagtgg ctccagttac 120
aaaaaaaaatt ttaatgaaaa cgttaaacat aaaaatagaa gtttgagatt ttaaaaaagtg 180
tataaaangc cccacaaaaac ttgtcaacgt tgttcccttat tctacaaaaat agcaccagta 240
agaagagtaa aaggtgttaa aaaccattat gacagcattt ctgaaatgca gcttgtctga 300

nttcccgttc tccctaaaaa cgacttctta tggataana aaggattaa aaaatctccn 360
aaaggaggc accgagcttt gcaggtttc cctggtcac tctcaggatg tggggggagg 420
gtatgggaa atggtatggt ctggtccctg gactggctgg tcactgcctc tggggttng 480
gtaaaagggt g 491

<210> 73
<211> 443
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (9)..(9)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (11)..(11)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (23)..(24)
<223> May be any nucleic acid

<220>
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<222> (126)..(126)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (157)..(157)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (170)..(170)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (341)..(341)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (347)..(347)
<223> May be any nucleic acid

<220>

<221> Unsure
<222> (371)..(371)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (405)..(405)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (412)..(412)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (430)..(430)
<223> May be any nucleic acid

<400> 73
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accggncgtt ctcttgatg tttgagaatg ttgtagnat gaaggttgn gacaagaggg 180
acatctcacg gtccctggag tgtaatccag tcatgattga tgccatcaaa gttctgtctg 240
ctcacagggc ccgatacttc tggggcaacc tacccggat gaacaggatc tttggcttgc 300
ctgtgcacta cacagacgtg tcccaacatg gggccgtggg ngccgcncca ggaagcttgc 360
tggggaaaggt nctggggagc gttgccttgt tcatcccac acctntttcg gnccctattg 420
qaaggqattn attttqcca tq 443

<210> 74
<211> 273
<212> DNA
<213> Homo sapiens

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<400> 74  
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ctggatttagt tctgatccc ccacaaggag ccctcgaatt ggctaaagtg agaaaactggg 120  
cctgaagact ccgtaccctc tgccatcttgc ccgagggagt ctccttttag aaaacaatca 180  
aagggttatt gcatgagtctt ggatgaatcc cactctcagc tgtccacggg cccgaccacc 240  
tcatctaqcc cccttttgg cagggagaac ctg 273
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<210>. 75
<211> 250
<212> DNA

<213> Homo sapiens

<220>
<221> Unsure
<222> (26)..(27)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (33)..(33)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (188)..(188)
<223> May be any nucleic acid

<400> 75
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aaactgttga aacaaggcac agtttttcc ccatatttg tacgtcgtag ctccagttac 120
aaaaaaaaatt ttaatgaaaa cgttaaacat aaaaatagaa gtttgagatt ttaaaaagtg 180
tataaaangc cccacaaaac ttgtcaacgt tgttccttat tctacaaaat agcaccagta 240
agaagagataa 250

<210> 76
<211> 443
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (9)..(9)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (11)..(11)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (23)..(24)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (126)..(126)

<223> May be any nucleic acid

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<221> Unsure

<222> (157)..(157)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (170)..(170)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (341)..(341)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (347)..(347)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (371)..(371)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (405)..(405)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (412)..(412)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (430)..(430)

<223> May be any nucleic acid

<400> 76

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accggncgtt cttctggatg tttgagaatg ttgtagnat gaaggttgn gacaagaggg 180
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